

2-Methyl-3-(10H-phenothiazin-10-yl)-buta-1,3-diene-1,1,4,4-tetracarbonitrile

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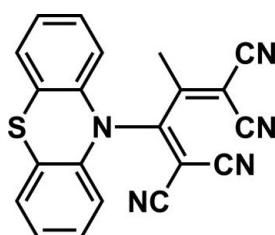
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Key indicators: single-crystal X-ray study; $T = 93\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.032; wR factor = 0.078; data-to-parameter ratio = 15.4.

In the title compound, $\text{C}_{21}\text{H}_{11}\text{N}_5\text{S}$, the phenothiazine unit has a butterfly structure, and the central six-membered ring adopts a boat conformation. The dihedral angle between the benzene rings is $127.64(6)^\circ$, which is smaller than those reported for similar compounds because of the steric repulsion between the phenothiazine and its tetracyano-1,3-butadiene substituent. The dicyanovinyl groups are almost orthogonal to one another, making a dihedral angle of $80.58(6)^\circ$. In the crystal, the molecules are aligned along the b axis. Four kinds of weak $\text{C}-\text{H}\cdots\text{N}$ interactions are recognized, one of which connects the molecules into a one-dimensional array and the remaining three link these arrays.

Related literature

For applications of tetracyano-1,3-butadienes in photonics and non-linear optics, see: Faupel *et al.* (2007). For the preparation and structure of 10-(prop-1-yn-1-yl)-10H-phenothiazine, see: Zaugg *et al.* (1958); Umezono & Okuno (2012). For the structures of other related *N*-substituted phenothiazines, see: Chu & Van der Helm (1974, 1975); Tokunaga & Okuno (2012).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{11}\text{N}_5\text{S}$

$M_r = 365.41$

Monoclinic, $P2_1$
 $a = 10.217(3)\text{ \AA}$
 $b = 7.848(3)\text{ \AA}$
 $c = 11.369(3)\text{ \AA}$
 $\beta = 97.316(4)^\circ$
 $V = 904.2(5)\text{ \AA}^3$

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.19\text{ mm}^{-1}$
 $T = 93\text{ K}$
 $0.10 \times 0.10 \times 0.05\text{ mm}$

Data collection

Rigaku Saturn724+ diffractometer
7524 measured reflections
3753 independent reflections
3494 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.078$
 $S = 1.04$
3753 reflections
244 parameters
1 restraint
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.21\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.25\text{ e \AA}^{-3}$
Absolute structure: Flack (1983),
1526 Friedel pairs
Flack parameter: 0.01 (6)

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C15—H15B···N4 ⁱ	0.98	2.65	3.186 (3)	114
C2—H2···N5 ⁱⁱ	0.95	2.59	3.352 (3)	137
C10—H10···N2 ⁱⁱⁱ	0.95	2.70	3.413 (3)	133
C8—H8···N2 ^{iv}	0.95	2.62	3.480 (3)	151

Symmetry codes: (i) $x, y + 1, z$; (ii) $-x, y - \frac{1}{2}, -z + 1$; (iii) $-x + 1, y - \frac{1}{2}, -z$; (iv) $-x + 1, y + \frac{1}{2}, -z$.

Data collection: *CrystalClear* (Rigaku, 2008); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *CrystalStructure* (Rigaku, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ5308).

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supplementary materials

Acta Cryst. (2013). E69, o665 [doi:10.1107/S1600536813008799]

2-Methyl-3-(10*H*-phenothiazin-10-yl)buta-1,3-diene-1,1,4,4-tetracarbonitrile

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Comment

Tetracyano-1,3-butadienes, which are prepared by TCNE addition to acetylene compounds carrying an electron donating part, have been attracted interest from the viewpoint of applications in photonics and nonlinear optics (Faupel *et al.*, 2007). The title compound, $C_{21}H_{11}N_5S_1$, is a TCNE adduct of 10-(prop-1-yn-1-yl)-10*H*-phenothiazine which is known as the first ynamine compound (Zaugg *et al.*, 1958).

The phenothiazine moiety has a butterfly structure, and the central six-membered ring adopts a boat conformation (Fig. 1). The dihedral angle between the C1—C6 and C7—C12 planes is $127.64(6)^\circ$. This is smaller than angles reported for related phenothiazine systems (Chu & Helm, 1974, 1975; Umezono & Okuno, 2012; Tokunaga & Okuno, 2012), and is presumably because of the steric repulsion between the phenothiazine ring system and its tetracyano-1,3-butadiene substituent. The structure around the N1 is pyrimidal and the N1 atom lies $0.1440(14)$ Å out of the C1/C12/C13 plane. This plane is inclined at $10.84(4)^\circ$ to the C13/C16—C18/N2/N3 dicyanovinyl plane (r.m.s. deviation = 0.0095 Å), indicating good π -conjugation. This latter plane is approximately orthogonal to the other C14/C19—C21/N4/N5 plane (r.m.s. deviation = 0.0170 Å), with a dihedral angle of $80.58(6)^\circ$ between them.

The molecules align along the *b* axis. Four kinds of weak C—H···N interactions are recognized in each molecule, one of which connects the molecules within the one-dimensional array and the remaining three link these arrays (Fig. 2, Table 1).

Experimental

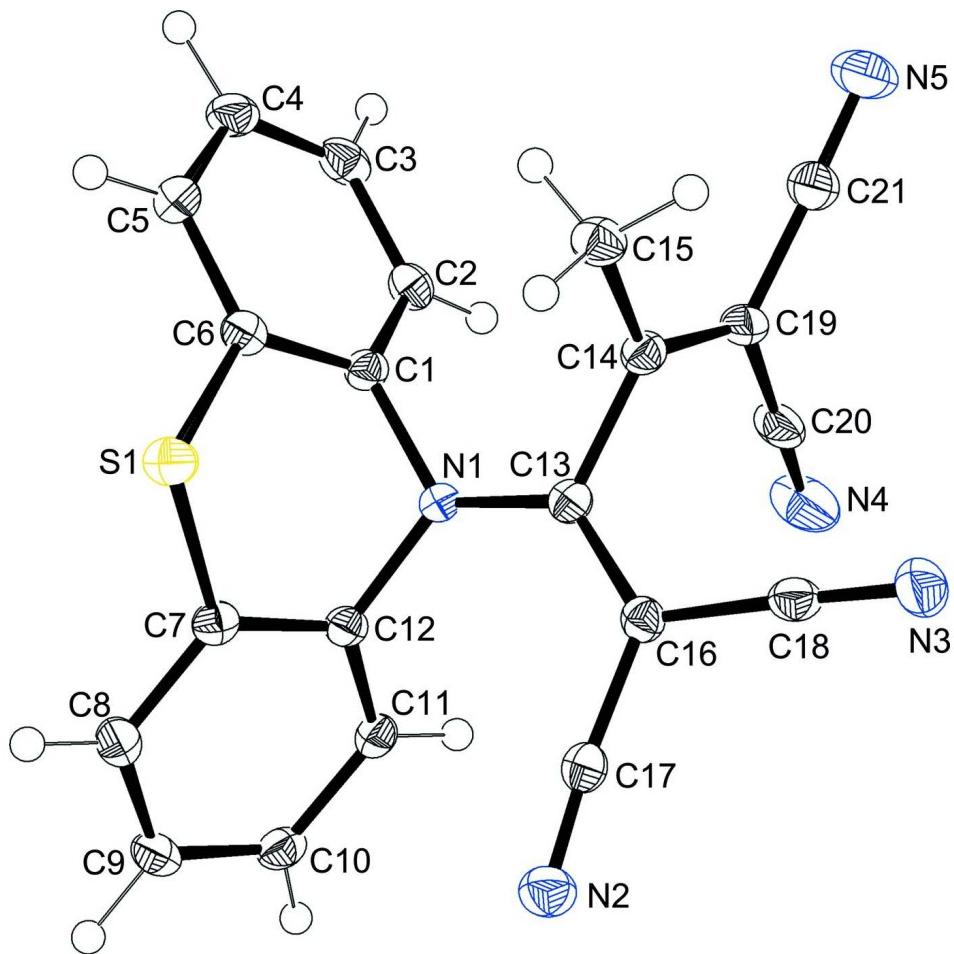
A solution of 10-(prop-1-yn-1-yl)-10*H*-phenothiazine (0.50 g, 2.1 mmol) (Zaugg *et al.*, 1958) in dichloromethane (25 ml) was added to a solution of ethene-1,1,2,2-tetracarbonitrile (0.27 g, 2.1 mmol) in dichloromethane (150 ml). The solution was stirred for a day and was concentrated by evaporation. The residue was purified by gel permeation chromatography (GPC) to give the title compound as pale blue powder (0.48 g, 62%). Single crystals with sufficient quality for X-ray crystallographic analysis were prepared by recrystallization from an acetonitrile solution.

Refinement

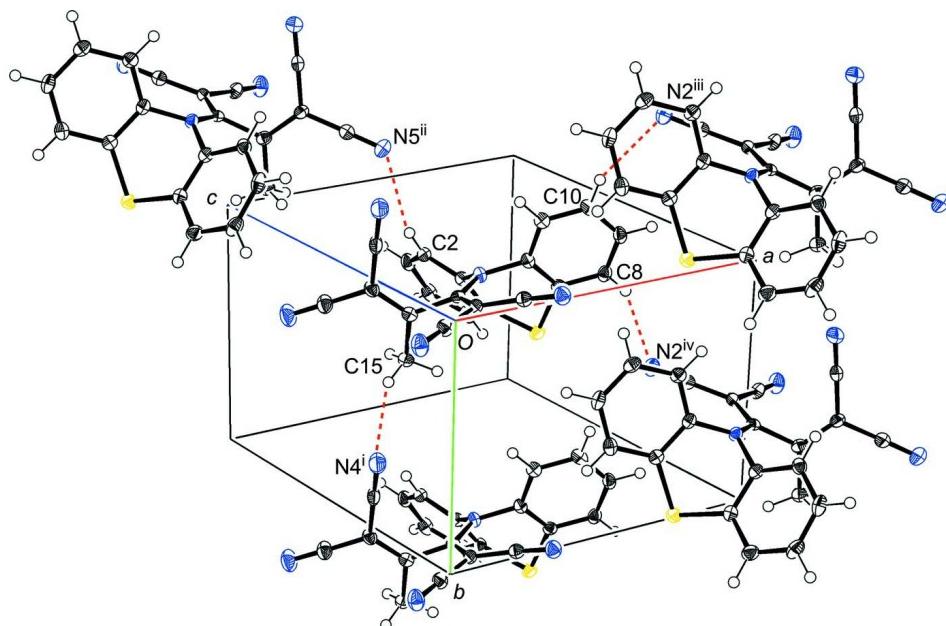
The C-bound H atoms were placed at ideal positions and were refined as riding on their parent C atoms. $U_{\text{iso}}(\text{H})$ values of the H atoms were set at $1.2U_{\text{eq}}(\text{parent atom})$.

Computing details

Data collection: *CrystalClear* (Rigaku, 2008); cell refinement: *CrystalClear* (Rigaku, 2008); data reduction: *CrystalClear* (Rigaku, 2008); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *CrystalStructure* (Rigaku, 2010).

**Figure 1**

The asymmetric unit of the title compound with atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres.

**Figure 2**

A view of the intermolecular interactions of the title compound. [Symmetry codes: (i) $x, y + 1, z$ (ii) $-x, y - 1/2, -z + 1$ (iii) $-x + 1, y - 1/2, -z$ (iv) $-x + 1, y + 1/2, -z$.]

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$C_{21}H_{11}N_3S$
 $M_r = 365.41$
Monoclinic, $P2_1$
Hall symbol: P 2yb
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 $b = 7.848 (3)$ Å
 $c = 11.369 (3)$ Å
 $\beta = 97.316 (4)^\circ$
 $V = 904.2 (5)$ Å³
 $Z = 2$

$F(000) = 376.00$
 $D_x = 1.342 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71075$ Å
Cell parameters from 3683 reflections
 $\theta = 1.8\text{--}30.7^\circ$
 $\mu = 0.19 \text{ mm}^{-1}$
 $T = 93 \text{ K}$
Block, pale blue
 $0.10 \times 0.10 \times 0.05$ mm

Data collection

Rigaku Saturn724+
diffractometer
Detector resolution: 28.445 pixels mm⁻¹
 ω scans
7524 measured reflections
3753 independent reflections

3494 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.022$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -12 \rightarrow 13$
 $k = -9 \rightarrow 10$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.078$
 $S = 1.04$
3753 reflections
244 parameters
1 restraint

Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.045P)^2 + 0.1327P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.21 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.25 \text{ e \AA}^{-3}$$

Absolute structure: Flack (1983), 1526 Friedel pairs

Flack parameter: 0.01 (6)

Special details

Refinement. Refinement was performed using all reflections. The weighted *R*-factor (*wR*) and goodness of fit (*S*) are based on F^2 . *R*-factor (gt) are based on *F*. The threshold expression of $F^2 > 2.0 \sigma(F^2)$ is used only for calculating *R*-factor (gt).

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.46875 (4)	0.28150 (5)	0.24265 (4)	0.02108 (10)
N1	0.27521 (13)	0.00720 (18)	0.24818 (12)	0.0165 (3)
N2	0.25743 (15)	-0.0784 (3)	-0.09700 (14)	0.0250 (4)
N3	-0.12618 (16)	0.0506 (3)	-0.02155 (14)	0.0300 (4)
N4	-0.03764 (19)	-0.3121 (3)	0.2637 (2)	0.0404 (5)
N5	-0.20384 (18)	0.1361 (3)	0.42695 (17)	0.0368 (5)
C1	0.30948 (16)	0.0842 (3)	0.36326 (14)	0.0173 (4)
C2	0.26062 (16)	0.0219 (3)	0.46368 (14)	0.0192 (4)
C3	0.29033 (17)	0.1081 (3)	0.57034 (15)	0.0239 (4)
C4	0.36780 (17)	0.2539 (3)	0.57534 (15)	0.0253 (4)
C5	0.42255 (17)	0.3100 (3)	0.47617 (16)	0.0235 (4)
C6	0.39519 (16)	0.2229 (3)	0.36953 (14)	0.0181 (4)
C7	0.48704 (16)	0.0732 (3)	0.18623 (13)	0.0167 (4)
C8	0.59958 (17)	0.0265 (3)	0.13638 (14)	0.0199 (4)
C9	0.61205 (17)	-0.1403 (3)	0.09858 (15)	0.0209 (4)
C10	0.51615 (17)	-0.2620 (3)	0.11218 (15)	0.0210 (4)
C11	0.40235 (15)	-0.2157 (3)	0.16050 (13)	0.0190 (3)
C12	0.38819 (16)	-0.0473 (3)	0.19417 (14)	0.0165 (4)
C13	0.15512 (16)	0.0349 (2)	0.18619 (14)	0.0162 (4)
C14	0.05534 (16)	0.1133 (3)	0.25659 (14)	0.0180 (4)
C15	0.05982 (17)	0.3019 (3)	0.27492 (15)	0.0218 (4)
C16	0.11722 (16)	0.0065 (3)	0.06727 (15)	0.0169 (4)
C17	0.19996 (17)	-0.0424 (3)	-0.01964 (15)	0.0190 (4)
C18	-0.01790 (17)	0.0327 (3)	0.01951 (14)	0.0208 (4)
C19	-0.03508 (17)	0.0124 (3)	0.29768 (15)	0.0202 (4)
C20	-0.03738 (18)	-0.1681 (3)	0.27904 (18)	0.0253 (4)
C21	-0.13076 (18)	0.0814 (3)	0.36816 (17)	0.0260 (4)
H2	0.2078	-0.0780	0.4593	0.0231*
H3	0.2577	0.0674	0.6398	0.0287*
H4	0.3837	0.3162	0.6474	0.0303*
H5	0.4783	0.4073	0.4814	0.0282*
H8	0.6668	0.1077	0.1284	0.0239*
H9	0.6875	-0.1719	0.0627	0.0250*
H10	0.5281	-0.3765	0.0886	0.0252*
H11	0.3361	-0.2976	0.1702	0.0228*
H15A	0.0728	0.3268	0.3601	0.0261*
H15B	-0.0234	0.3525	0.2389	0.0261*

H15C	0.1330	0.3500	0.2378	0.0261*
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Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0262 (2)	0.0173 (2)	0.0209 (2)	-0.00254 (18)	0.00752 (15)	0.00042 (16)
N1	0.0147 (7)	0.0192 (8)	0.0154 (7)	-0.0000 (6)	0.0014 (5)	-0.0034 (6)
N2	0.0252 (8)	0.0294 (9)	0.0208 (8)	0.0023 (7)	0.0040 (6)	-0.0002 (7)
N3	0.0214 (8)	0.0444 (11)	0.0240 (8)	0.0023 (8)	0.0017 (7)	-0.0006 (7)
N4	0.0411 (11)	0.0224 (10)	0.0632 (14)	-0.0058 (8)	0.0282 (10)	-0.0079 (9)
N5	0.0355 (10)	0.0319 (11)	0.0470 (11)	-0.0014 (8)	0.0204 (9)	-0.0128 (8)
C1	0.0180 (8)	0.0184 (9)	0.0151 (8)	0.0022 (7)	0.0008 (6)	-0.0008 (7)
C2	0.0175 (8)	0.0192 (9)	0.0215 (8)	0.0015 (7)	0.0040 (7)	0.0028 (7)
C3	0.0217 (9)	0.0338 (11)	0.0168 (8)	0.0078 (8)	0.0047 (7)	0.0035 (7)
C4	0.0232 (9)	0.0339 (12)	0.0180 (8)	0.0067 (8)	-0.0003 (7)	-0.0069 (8)
C5	0.0203 (8)	0.0246 (11)	0.0251 (9)	-0.0011 (8)	0.0009 (7)	-0.0074 (7)
C6	0.0180 (8)	0.0194 (8)	0.0171 (8)	0.0018 (7)	0.0030 (7)	0.0009 (6)
C7	0.0187 (8)	0.0174 (8)	0.0136 (7)	0.0018 (7)	0.0002 (6)	0.0012 (6)
C8	0.0175 (8)	0.0266 (9)	0.0154 (8)	0.0004 (8)	0.0018 (6)	0.0037 (7)
C9	0.0183 (8)	0.0294 (10)	0.0152 (8)	0.0064 (8)	0.0034 (6)	0.0018 (7)
C10	0.0221 (9)	0.0224 (10)	0.0172 (8)	0.0048 (7)	-0.0020 (7)	-0.0039 (6)
C11	0.0177 (8)	0.0210 (8)	0.0175 (7)	0.0000 (8)	-0.0009 (6)	-0.0019 (7)
C12	0.0156 (8)	0.0210 (8)	0.0126 (7)	0.0030 (7)	0.0002 (6)	-0.0011 (6)
C13	0.0180 (8)	0.0109 (8)	0.0199 (8)	-0.0016 (7)	0.0033 (6)	-0.0001 (6)
C14	0.0188 (9)	0.0173 (9)	0.0173 (8)	0.0015 (7)	-0.0001 (6)	-0.0016 (6)
C15	0.0243 (9)	0.0173 (10)	0.0244 (8)	0.0010 (8)	0.0057 (7)	-0.0003 (7)
C16	0.0159 (8)	0.0155 (8)	0.0195 (8)	-0.0017 (7)	0.0033 (6)	0.0002 (7)
C17	0.0184 (8)	0.0193 (9)	0.0186 (8)	-0.0000 (7)	0.0003 (7)	0.0007 (6)
C18	0.0230 (10)	0.0238 (9)	0.0159 (8)	0.0027 (8)	0.0037 (7)	-0.0016 (7)
C19	0.0197 (8)	0.0179 (9)	0.0234 (8)	0.0010 (7)	0.0045 (7)	-0.0050 (7)
C20	0.0208 (9)	0.0246 (10)	0.0331 (10)	-0.0042 (7)	0.0139 (8)	-0.0049 (8)
C21	0.0234 (10)	0.0231 (10)	0.0329 (10)	-0.0047 (8)	0.0090 (8)	-0.0066 (8)

Geometric parameters (\AA , $^\circ$)

S1—C6	1.7701 (18)	C11—C12	1.389 (3)
S1—C7	1.7745 (19)	C13—C14	1.505 (3)
N1—C1	1.443 (2)	C13—C16	1.376 (3)
N1—C12	1.440 (3)	C14—C15	1.495 (3)
N1—C13	1.352 (2)	C14—C19	1.345 (3)
N2—C17	1.153 (3)	C16—C17	1.432 (3)
N3—C18	1.153 (3)	C16—C18	1.432 (3)
N4—C20	1.143 (3)	C19—C20	1.432 (3)
N5—C21	1.147 (3)	C19—C21	1.446 (3)
C1—C2	1.392 (3)	C2—H2	0.950
C1—C6	1.393 (3)	C3—H3	0.950
C2—C3	1.388 (3)	C4—H4	0.950
C3—C4	1.388 (3)	C5—H5	0.950
C4—C5	1.392 (3)	C8—H8	0.950
C5—C6	1.389 (3)	C9—H9	0.950

C7—C8	1.394 (3)	C10—H10	0.950
C7—C12	1.395 (3)	C11—H11	0.950
C8—C9	1.389 (3)	C15—H15A	0.980
C9—C10	1.391 (3)	C15—H15B	0.980
C10—C11	1.396 (3)	C15—H15C	0.980
C6—S1—C7	97.49 (9)	C13—C16—C18	119.09 (16)
C1—N1—C12	113.32 (13)	C17—C16—C18	113.70 (15)
C1—N1—C13	120.28 (14)	N2—C17—C16	174.02 (18)
C12—N1—C13	123.30 (14)	N3—C18—C16	178.0 (2)
N1—C1—C2	121.70 (15)	C14—C19—C20	122.01 (18)
N1—C1—C6	116.87 (15)	C14—C19—C21	120.99 (18)
C2—C1—C6	121.43 (15)	C20—C19—C21	116.93 (17)
C1—C2—C3	118.95 (17)	N4—C20—C19	179.1 (3)
C2—C3—C4	119.83 (17)	N5—C21—C19	178.0 (2)
C3—C4—C5	120.88 (17)	C1—C2—H2	120.520
C4—C5—C6	119.61 (17)	C3—C2—H2	120.526
S1—C6—C1	119.39 (13)	C2—C3—H3	120.088
S1—C6—C5	121.62 (14)	C4—C3—H3	120.079
C1—C6—C5	118.99 (16)	C3—C4—H4	119.564
S1—C7—C8	121.29 (14)	C5—C4—H4	119.554
S1—C7—C12	119.38 (13)	C4—C5—H5	120.189
C8—C7—C12	119.32 (16)	C6—C5—H5	120.199
C7—C8—C9	119.11 (17)	C7—C8—H8	120.448
C8—C9—C10	121.28 (17)	C9—C8—H8	120.446
C9—C10—C11	119.94 (18)	C8—C9—H9	119.360
C10—C11—C12	118.51 (17)	C10—C9—H9	119.363
N1—C12—C7	116.93 (15)	C9—C10—H10	120.029
N1—C12—C11	121.16 (15)	C11—C10—H10	120.035
C7—C12—C11	121.71 (16)	C10—C11—H11	120.744
N1—C13—C14	114.81 (14)	C12—C11—H11	120.746
N1—C13—C16	127.51 (16)	C14—C15—H15A	109.470
C14—C13—C16	117.63 (14)	C14—C15—H15B	109.468
C13—C14—C15	117.89 (15)	C14—C15—H15C	109.466
C13—C14—C19	119.08 (16)	H15A—C15—H15B	109.474
C15—C14—C19	123.01 (17)	H15A—C15—H15C	109.475
C13—C16—C17	127.17 (15)	H15B—C15—H15C	109.474

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C15—H15B···N4 ⁱ	0.98	2.65	3.186 (3)	114
C2—H2···N5 ⁱⁱ	0.95	2.59	3.352 (3)	137
C10—H10···N2 ⁱⁱⁱ	0.95	2.70	3.413 (3)	133
C8—H8···N2 ^{iv}	0.95	2.62	3.480 (3)	151

Symmetry codes: (i) $x, y+1, z$; (ii) $-x, y-1/2, -z+1$; (iii) $-x+1, y-1/2, -z$; (iv) $-x+1, y+1/2, -z$.